

Final TCR Display PCT.WorkFile.txt

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<110> FirstName : Bent
<110> MiddleInitial : K
<110> Suffix : Dr

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<110> FirstName : Torben
<110> MiddleInitial : B
<110> Suffix : Dr

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<110> LastName : Molloy
<110> FirstName : Peter
<110> MiddleInitial : E
<110> Suffix : Dr

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<110> LastName : Boulter

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<110> Suffix : Dr

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<110> LastName : Li
<110> FirstName : Yi
<110> MiddleInitial :
<110> Suffix : Dr

Application Project

<120> Title : Substances
<130> AppFileReference : Case No 19
<140> CurrentAppNumber : GB 0226227.7
<141> CurrentFilingDate : 2002-11-09

Earlier Applications

<150> PriorAppNumber : GB0226227.7
<151> PriorFilingDate : 2002-11-09

Earlier Applications

<150> PriorAppNumber : GB 0301814.0
<151> PriorFilingDate : 2003-01-25

Earlier Applications

<150> PriorAppNumber : GB 0304067.2
<151> PriorFilingDate : 2003-02-22

Earlier Applications

<150> PriorAppNumber : GB 0311397.4
<151> PriorFilingDate : 2003-05-16

Earlier Applications

<150> PriorAppNumber : GB 0316356.5
<151> PriorFilingDate : 2003-07-11

Earlier Applications

<150> PriorAppNumber : US 60/436,323
<151> PriorFilingDate : 2003-04-16

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
DSDVYITDKT VLDMRSMDFK
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 1
SequenceDescription :

Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
QSKDSDVYIT DKTVLDMRSM 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 2
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
DIQNPDPAVY QLRDSKSSDK 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 3
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
DPAVYQLRDS KSSDKSVCLF 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 4
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
NGKEVHSGVS TDPQLKEQP 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 5
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ALNDSRYALS SRLRVSATFW 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 6
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
PPEVAVFEPS EAEISHTQKA 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 7
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
KEVHSGVSTD PQPLKEQPAL 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 8
SequenceDescription :

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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
VFPPEVAVFE PSEAEISHTQ 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 9
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
PYIQNPEPAV YQLKDPRSQD STLCLFTDFD SQINVPKTME SGTFITDKTV LDMKAMDSKS 60
NGAIAWSNQT SFTCQDIFKE TNATYPSSDV P 91
<212> Type : PRT
<211> Length : 91
SequenceName : SEQ ID 10
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
EDLRNVTPPK VSLFEPSKAE IANKQKATLV CLARGFFPDH VELSWWVNGR EVHSGVSTDP 60
QAYKESNYSY CLSSRLRVSA TFWHNPRNHF RCQVQFHGLS EEDKWPEGSP KPVTQNISAE 120
AWGRAD 126
<212> Type : PRT
<211> Length : 126
SequenceName : SEQ ID 11
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ESGTFITDKT VLDMKAMDSK 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 12
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
KTMESGTFIT DKTVLDMKAM 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 13
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
YIQNPEPAVY QLKDPRSQDS 20
<212> Type : PRT
<211> Length : 20
SequenceName : SEQ ID 14
SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 AVYQLKDPRS QDSTLCLFTD 20
 <212> Type : PRT
 <211> Length : 20
 SequenceName : SEQ ID 15
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 NGREVHSGVS TDPQAYKESN 20
 <212> Type : PRT
 <211> Length : 20
 SequenceName : SEQ ID 16
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 KESNYSYCLS SRLRVSATFW 20
 <212> Type : PRT
 <211> Length : 20
 SequenceName : SEQ ID 17
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 PPKVSLFEPS KAEIANKQKA 20
 <212> Type : PRT
 <211> Length : 20
 SequenceName : SEQ ID 18
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 REVHSGVSTD PQAYKESNYS 20
 <212> Type : PRT
 <211> Length : 20
 SequenceName : SEQ ID 19
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 VTPPKVSLFE PSKAEIANKQ 20
 <212> Type : PRT
 <211> Length : 20
 SequenceName : SEQ ID 20
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 LLFGYPVYV 9
 <212> Type : PRT
 <211> Length : 9
 SequenceName : SEQ ID 21
 SequenceDescription :

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Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
SLLMITQC 8
<212> Type : PRT
<211> Length : 8
SequenceName : SEQ ID 22
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
LLGRNSFEV 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 23
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
KLVALGINAV 10
<212> Type : PRT
<211> Length : 10
SequenceName : SEQ ID 24
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
LLGDLFGV 8
<212> Type : PRT
<211> Length : 8
SequenceName : SEQ ID 25
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
FLRGRAYGL 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 26
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
HRCQAIRKK 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 27
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
YRSGIIAVV 9
<212> Type : PRT
<211> Length : 9
SequenceName : SEQ ID 28

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SequenceDescription :

Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
VYGFVRACL                                     9
<212> Type : PRT
<211> Length : 9
      SequenceName : SEQ ID 29
      SequenceDescription :
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Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ILAKFLHWL                                     9
<212> Type : PRT
<211> Length : 9
      SequenceName : SEQ ID 30
      SequenceDescription :
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Sequence

```
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
LTLGEFLKL                                     9
<212> Type : PRT
<211> Length : 9
      SequenceName : SEQ ID 31
      SequenceDescription :
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Sequence

```
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
PKYVKQNTLK LA                                12
<212> Type : PRT
<211> Length : 12
      SequenceName : SEQ ID 32
      SequenceDescription :
```

Sequence

```
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
GILGFVFTL                                     9
<212> Type : PRT
<211> Length : 9
      SequenceName : SEQ ID 33
      SequenceDescription :
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Sequence

```
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
SLYNTVATL                                     9
<212> Type : PRT
<211> Length : 9
      SequenceName : SEQ ID 34
      SequenceDescription :
```

Sequence

```
-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cacagacaaa tgtgtgctag acat                   24
<212> Type : DNA
```

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<211> Length : 24
SequenceName : SEQ ID 35
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 35

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atgtctagca cacattgtc tgtg
<212> Type : DNA
<211> Length : 24
SequenceName : SEQ ID 36
SequenceDescription :

24

Custom Codon

Sequence Name : SEQ ID 36

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cagtggggtc tgcacagacc c
<212> Type : DNA
<211> Length : 21
SequenceName : SEQ ID 37
SequenceDescription :

21

Custom Codon

Sequence Name : SEQ ID 37

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gggtctgtgc agacccact g
<212> Type : DNA
<211> Length : 21
SequenceName : SEQ ID 38
SequenceDescription :

21

Custom Codon

Sequence Name : SEQ ID 38

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
taataatagc tataataata ttctatttca aggagacagt c
<212> Type : DNA
<211> Length : 41
SequenceName : SEQ ID 39
SequenceDescription :

41

Custom Codon

Sequence Name : SEQ ID 39

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :

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caatccagcg gctgccgtag gcaataggta ttccattatg actgtctcct tgaaatag 58

<212> Type : DNA
<211> Length : 58
SequenceName : SEQ ID 40
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 40

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctacggcagc cgctggattg ttattactcg cggcccagcc ggccatggcc cag 53

<212> Type : DNA
<211> Length : 53
SequenceName : SEQ ID 41
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 41

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gttctgctcc acttccttct gggccatggc cggtggggc g 41
<212> Type : DNA
<211> Length : 41
SequenceName : SEQ ID 42
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 42

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cagaaggaag tggagcagaa c 21
<212> Type : DNA
<211> Length : 21
SequenceName : SEQ ID 43
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 43

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cttcttaaag aattcttaat taacctaggt tattaggaac tttctgggct ggggaag 57

<212> Type : DNA
<211> Length : 57
SequenceName : SEQ ID 44
SequenceDescription :

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Custom Codon

Sequence Name : SEQ ID 44

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gttaattaag aattctttaa gaaggagata tacatatgaa aaaattatta ttcgcaattc 60<212> Type : DNA
<211> Length : 60
SequenceName : SEQ ID 45
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 45

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cgcgctgtga gaatagaaag gaacaactaa aggaattgcg aataataatt ttttcatatg 60<212> Type : DNA
<211> Length : 60
SequenceName : SEQ ID 46
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 46

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctttctattc tcacagcgcg caggctggtg tcactcagac 40
<212> Type : DNA
<211> Length : 40
SequenceName : SEQ ID 47
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 47

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atgatgtcta gatgcggccg cgtctgctct accccaggcc tc 42
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 48
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 48

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :

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gcatctagac atcatcacca tcatcactag actgttgaaa gttgttttagc aaaac 55

<212> Type : DNA
<211> Length : 55
SequenceName : SEQ ID 49
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 49

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctagagggtta cttattaag actccttatt acgcagtatg 40
<212> Type : DNA
<211> Length : 40
SequenceName : SEQ ID 50
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 50

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
agctgcagct aatacgactc actataggaa caggccacca tggcgtcgat tatgctgagt 60
gatatccttg tccggtggta ccctag 86
<212> Type : DNA
<211> Length : 86
SequenceName : SEQ ID 51
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 51

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gatcccatgg tggcctgttc ctatagttagc tcgtattagc tgc 43
<212> Type : DNA
<211> Length : 43
SequenceName : SEQ ID 52
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 52

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
agctgcagct aatacgactc actataggaa caggccacca tgg 43
<212> Type : DNA
<211> Length : 43
SequenceName : SEQ ID 53
SequenceDescription :

Custom Codon

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Sequence Name : SEQ ID 53

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ccaccatggg ccagaaggaa gtggagcaga actc

34

<212> Type : DNA

<211> Length : 34

SequenceName : SEQ ID 54

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 54

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

cgagagcccg tagaactgga cttg

24

<212> Type : DNA

<211> Length : 24

SequenceName : SEQ ID 55

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 55

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gtggatccgg cggtggcggg tcgaacgctg gtgtcactca gacccc

46

<212> Type : DNA

<211> Length : 46

SequenceName : SEQ ID 56

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 56

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ccggatccac ctccgcctga accgcctcca ccggtgacca caacctgggt ccctg

55

<212> Type : DNA

<211> Length : 55

SequenceName : SEQ ID 57

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 57

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ctgagaattc ttatgactct ccgcggttga agctc

35

<212> Type : DNA

<211> Length : 35

SequenceName : SEQ ID 58

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SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 58

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

tgacgaattc tgactctccg cggttgaagc tc

32

<212> Type : DNA

<211> Length : 32

SequenceName : SEQ ID 59

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 59

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

agctgcagct aatacgaactc actatagg

28

<212> Type : DNA

<211> Length : 28

SequenceName : SEQ ID 60

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 60

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ggccaccatg ggcaacgctg gtgtcactca gacccc

36

<212> Type : DNA

<211> Length : 36

SequenceName : SEQ ID 61

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 61

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

tgaaccgcct ccaccgtctg ctctacccca ggcctcggcg

40

<212> Type : DNA

<211> Length : 40

SequenceName : SEQ ID 62

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 62

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

tgactctccg cggttgaagc tc

22

<212> Type : DNA

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<211> Length : 22
SequenceName : SEQ ID 63
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 63

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cagctggggg aagcttcagt ttggagcag
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 64
SequenceDescription :

29

Custom Codon

Sequence Name : SEQ ID 64

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctgctccaaa ctgaagcttc ccccagctg
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 65
SequenceDescription :

29

Custom Codon

Sequence Name : SEQ ID 65

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gtacttctgt gcctcgaggc cgggactag
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 66
SequenceDescription :

29

Custom Codon

Sequence Name : SEQ ID 66

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctagtcccg cctcgaggca cagaagtac
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 67
SequenceDescription :

29

Custom Codon

Sequence Name : SEQ ID 67

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :

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catttttcagg gatagcaagc 20
<212> Type : DNA
<211> Length : 20
SequenceName : SEQ ID 68
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 68

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString : 40
tcacacagga aacagctatg tcacacagga aacagctatg
<212> Type : DNA
<211> Length : 40
SequenceName : SEQ ID 69
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 69

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString : 49
tgtgcctcga ggnnknnknn knnknnknnk cgaccagagc agtacttcg
<212> Type : DNA
<211> Length : 49
SequenceName : SEQ ID 70
SequenceDescription :

Feature

Sequence: SEQ ID 70:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 49
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 70

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString : 52
tgtgcctcga ggccgnnknn knnknnknnk nnkccagagc agtacttcgg gc

<212> Type : DNA
<211> Length : 52
SequenceName : SEQ ID 71
SequenceDescription :

Feature

Sequence: SEQ ID 71:
<221> FeatureKey : misc_feature
<222> LocationFrom :
<222> LocationTo :
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

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Feature

Sequence: SEQ ID 71:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 52
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 71

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggccggnknkn knknknknkn nnkcgaccag agcagtactt cg 52

<212> Type : DNA
<211> Length : 52
SequenceName : SEQ ID 72
SequenceDescription :

Feature

Sequence: SEQ ID 72:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 52
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 72

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggccggnknkn knknknknkn nnkggagggc gaccagagca g 51

<212> Type : DNA
<211> Length : 51
SequenceName : SEQ ID 73
SequenceDescription :

Feature

Sequence: SEQ ID 73:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 51
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 73

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggccgggann knknknknkn nnknknkggc gaccagagca gtac 54

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<212> Type : DNA
<211> Length : 54
SequenceName : SEQ ID 74
SequenceDescription :

Feature

Sequence: SEQ ID 74:
<221> FeatureKey : misc_feature
<222> LocationFrom :
<222> LocationTo :
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Feature

Sequence: SEQ ID 74:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 54
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 74

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggnnknnknn knnknnknnk ccagagcagt acttcgggc 49
<212> Type : DNA
<211> Length : 49
SequenceName : SEQ ID 75
SequenceDescription :

Feature

Sequence: SEQ ID 75:
<221> FeatureKey : misc_feature
<222> LocationFrom :
<222> LocationTo :
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Feature

Sequence: SEQ ID 75:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 49
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 75

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tgtgcctcga ggnnknnknn knnknnknnk gagcagtact tcgggccg 48
<212> Type : DNA
<211> Length : 48

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SequenceName : SEQ ID 76

SequenceDescription :

Feature

Sequence: SEQ ID 76:

<221> FeatureKey : misc_feature

<222> LocationFrom : 1

<222> LocationTo : 48

Other Information : m is a or c; n is a or t or g or c; k is g or t

CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 76

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

tgtgcctcga ggnnknnknn knnknnknnk cagtacttcg ggccgggc

48

<212> Type : DNA

<211> Length : 48

SequenceName : SEQ ID 77

SequenceDescription :

Feature

Sequence: SEQ ID 77:

<221> FeatureKey : misc_feature

<222> LocationFrom : 1

<222> LocationTo : 48

Other Information : m is a or c; n is a or t or g or c; k is g or t

CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 77

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

tgtgcctcga ggccgnnknn knnknnkggg cgaccagagc agtacttcg

49

<212> Type : DNA

<211> Length : 49

SequenceName : SEQ ID 78

SequenceDescription :

Feature

Sequence: SEQ ID 78:

<221> FeatureKey : misc_feature

<222> LocationFrom :

<222> LocationTo :

Other Information : m is a or c; n is a or t or g or c; k is g or t

CDSJoin : No

Feature

Sequence: SEQ ID 78:

<221> FeatureKey : misc_feature

<222> LocationFrom : 1

<222> LocationTo : 49

Other Information : m is a or c; n is a or t or g or c; k is g or t

CDSJoin : No

Custom Codon

Final TCR Display PCT.WorkFile.txt

Sequence Name : SEQ ID 78

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
aaactgaagc ttmnnmnnmn nmnnmntgt aacggcacag aggtag 46
<212> Type : DNA
<211> Length : 46
SequenceName : SEQ ID 79
SequenceDescription :

Feature

Sequence: SEQ ID 79:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 46
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 79

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
aaactgaagc ttmnnmnnngc tgctmnnntgt aacggcacag aggtag 46
<212> Type : DNA
<211> Length : 46
SequenceName : SEQ ID 80
SequenceDescription :

Feature

Sequence: SEQ ID 80:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 46
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 80

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
aaactgaagc ttmnnmnnmn ngctgtcmnn tgtaacggca cagaggtag 49
<212> Type : DNA
<211> Length : 49
SequenceName : SEQ ID 81
SequenceDescription :

Feature

Sequence: SEQ ID 81:
<221> FeatureKey : misc_feature
<222> LocationFrom :
<222> LocationTo :
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Final TCR Display PCT.WorkFile.txt

Feature

Sequence: SEQ ID 81:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 49
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 81

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
aaactgaagc ttmnnmnngc tgctcmnaac ggcacagagg tag 43
<212> Type : DNA
<211> Length : 43
SequenceName : SEQ ID 82
SequenceDescription :

Feature

Sequence: SEQ ID 82:
<221> FeatureKey : misc_feature
<222> LocationFrom : 1
<222> LocationTo : 43
Other Information : m is a or c; n is a or t or g or c; k is g or t
CDSJoin : No

Custom Codon

Sequence Name : SEQ ID 82

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggactagc gggagggcga ccagagcagt ag 42
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 83
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 83

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLAGGR PEQY 14
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 84
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cggggctgat gaggcgtag ccagagcagt ac 42
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 85

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SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 85

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 ASRPGLMSAX PEQY 14
 <212> Type : PRT
 <211> Length : 14
 SequenceName : SEQ ID 86
 SequenceDescription :

Feature

Sequence: SEQ ID 86:
 <221> FeatureKey : MISC_FEATURE
 <222> LocationFrom : 1
 <222> LocationTo : 14
 Other Information : x denotes an amino acid residue encoded by an amber
 codon
 CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gcctcgaggc cggggctgag gtcggcgtag ccagagcagt ac 42
 <212> Type : DNA
 <211> Length : 42
 SequenceName : SEQ ID 87
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 87

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 ASRPGLRSAX PEQY 14
 <212> Type : PRT
 <211> Length : 14
 SequenceName : SEQ ID 88
 SequenceDescription :

Feature

Sequence: SEQ ID 88:
 <221> FeatureKey : MISC_FEATURE
 <222> LocationFrom : 1
 <222> LocationTo : 14
 Other Information : x denotes an amino acid residue encoded by an amber
 codon
 CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 gcctcgaggc cgggactagc gggagggcga ccagagggcgt ag 42
 <212> Type : DNA
 <211> Length : 42
 SequenceName : SEQ ID 89

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SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 89

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ASRPGLAGGR PEAX

14

<212> Type : PRT

<211> Length : 14

SequenceName : SEQ ID 90

SequenceDescription :

Feature

Sequence: SEQ ID 90:

<221> FeatureKey : MISC_FEATURE

<222> LocationFrom :

<222> LocationTo :

Other Information : x denotes an amino acid residue encoded by an amber codon

CDSJoin : No

Feature

Sequence: SEQ ID 90:

<221> FeatureKey : MISC_FEATURE

<222> LocationFrom : 1

<222> LocationTo : 14

Other Information : x denotes an amino acid residue encoded by an amber codon

CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gcctcgaggc cgggactagc gggagggcgga ccagaggatt ag

42

<212> Type : DNA

<211> Length : 42

SequenceName : SEQ ID 91

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 91

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ASRPGLAGGR PEDX

14

<212> Type : PRT

<211> Length : 14

SequenceName : SEQ ID 92

SequenceDescription :

Feature

Sequence: SEQ ID 92:

<221> FeatureKey : MISC_FEATURE

<222> LocationFrom : 1

<222> LocationTo : 14

Other Information : x denotes an amino acid residue encoded by an amber codon

Final TCR Display PCT.WorkFile.txt

CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggactagc gggagggcga ccagatcagt ag
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 93
SequenceDescription :

42

Custom Codon

Sequence Name : SEQ ID 93

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLAGGR PDQX
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 94
SequenceDescription :

14

Feature

Sequence: SEQ ID 94:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
Other Information : x denotes an amino acid residue encoded by an amber
codon
CDSJoin : No

Feature

Sequence: SEQ ID 94:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber
codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggtctgta ggctgggcga ccagagcagt ac
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 95
SequenceDescription :

42

Custom Codon

Sequence Name : SEQ ID 95

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLXAGR PEQY
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 96

14

Final TCR Display PCT.WorkFile.txt
SequenceDescription :

Feature

Sequence: SEQ ID 96:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
Other Information : x denotes an amino acid residue encoded by an amber
codon
CDSJoin : No

Feature

Sequence: SEQ ID 96:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber
codon
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cggggctggt tccggggcga ccagagcagt ag 42
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 97
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 97

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLVPGR PEQX 14
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 98
SequenceDescription :

Feature

Sequence: SEQ ID 98:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
Other Information : x denotes an amino acid residue encoded by an amber
codon
CDSJoin : No

Feature

Sequence: SEQ ID 98:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber
codon
CDSJoin : No

Sequence

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```

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cggggcttgt gtctgcttag ccagagcagt ac
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 99
SequenceDescription :

```

Custom Codon

Sequence Name : SEQ ID 99

Sequence

```

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLVSAX PEQY
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 100
SequenceDescription :

```

Feature

```

Sequence: SEQ ID 100:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 14
Other Information : x denotes an amino acid residue encoded by an amber
codon
CDSJoin : No

```

Sequence

```

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gcctcgaggc cgggactagc gggagggcga ccacatccgt ag
<212> Type : DNA
<211> Length : 42
SequenceName : SEQ ID 101
SequenceDescription :

```

Custom Codon

Sequence Name : SEQ ID 101

Sequence

```

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ASRPGLAGGR PHPX
<212> Type : PRT
<211> Length : 14
SequenceName : SEQ ID 102
SequenceDescription :

```

Feature

```

Sequence: SEQ ID 102:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
Other Information : x denotes an amino acid residue encoded by an amber
codon
CDSJoin : No

```

Feature

Final TCR Display PCT.WorkFile.txt

Sequence: SEQ ID 102:

<221> FeatureKey : MISC_FEATURE

<222> LocationFrom : 1

<222> LocationTo : 14

Other Information : x denotes an amino acid residue encoded by an amber codon

CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gcctcgaggc cgggactagc gggagggcga ccagatgcgt ag 42

<212> Type : DNA

<211> Length : 42

SequenceName : SEQ ID 103

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 103

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ASRPGLAGGR PDAX 14

<212> Type : PRT

<211> Length : 14

SequenceName : SEQ 104

SequenceDescription :

Feature

Sequence: SEQ 104:

<221> FeatureKey : MISC_FEATURE

<222> LocationFrom : 1

<222> LocationTo : 14

Other Information : x denotes an amino acid residue encoded by an amber

codon

CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gcctcgaggc cgggtctgat tagtgcttag ccagagcagt ac 42

<212> Type : DNA

<211> Length : 42

SequenceName : SEQ ID 105

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 105

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ASRPGLISAX PEQY 14

<212> Type : PRT

<211> Length : 14

SequenceName : SEQ ID 106

SequenceDescription :

Feature

Final TCR Display PCT.WorkFile.txt

Sequence: SEQ ID 106:

<221> FeatureKey : MISC_FEATURE

<222> LocationFrom : 1

<222> LocationTo : 14

Other Information : x denotes an amino acid residue encoded by an amber codon

CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gccgttacaa ctgacagctg ggggaagctt cag 33

<212> Type : DNA

<211> Length : 33

SequenceName : SEQ ID 107

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 107

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

AVTTDSWGKL Q 11

<212> Type : PRT

<211> Length : 11

SequenceName : SEQ ID 108

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gccgttacaa ctgacagctg ggggccgctt cag 33

<212> Type : DNA

<211> Length : 33

SequenceName : SEQ ID 109

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 109

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

AVTTDSWGPL Q 11

<212> Type : PRT

<211> Length : 11

SequenceName : SEQ ID 110

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gccgttacaa ctgacagctg ggggaagatg cag 33

<212> Type : DNA

<211> Length : 33

SequenceName : SEQ ID 111

SequenceDescription :

Custom Codon

Final TCR Display PCT.WorkFile.txt

Sequence Name : SEQ ID 111

Sequence

```
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKM Q 11
<212> Type : PRT
<211> Length : 11
      SequenceName : SEQ ID 112
      SequenceDescription :
```

Sequence

```
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccggttaca ctgacagctg ggggaagttg cat 33
<212> Type : DNA
<211> Length : 33
      SequenceName : SEQ ID 113
      SequenceDescription :
```

Custom Codon

Sequence Name : SEQ ID 113

Sequence

```
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKL H 11
<212> Type : PRT
<211> Length : 11
      SequenceName : SEQ ID 114
      SequenceDescription :
```

Sequence

```
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccggttaca ctgacagctg ggggtagctt cat 33
<212> Type : DNA
<211> Length : 33
      SequenceName : SEQ ID 115
      SequenceDescription :
```

Custom Codon

Sequence Name : SEQ ID 115

Sequence

```
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGXL H 11
<212> Type : PRT
<211> Length : 11
      SequenceName : SEQ ID 116
      SequenceDescription :
```

Feature

```
Sequence: SEQ ID 116:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom :
<222> LocationTo :
      Other Information : x denotes an amino acid residue encoded by an amber
codon
```

Final TCR Display PCT.WorkFile.txt

CDSJoin : No

Feature

Sequence: SEQ ID 116:

<221> FeatureKey : MISC_FEATURE

<222> LocationFrom : 1

<222> LocationTo : 11

Other Information : x denotes an amino acid residue encoded by an amber codon

CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gccggttaca ctgacagctg gggggagctt cat

33

<212> Type : DNA

<211> Length : 33

SequenceName : SEQ ID 117

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 117

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

AVTTDSWGEL H

11

<212> Type : PRT

<211> Length : 11

SequenceName : SEQ ID 118

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gccggttaca ctgacagctg ggggaggctg cat

33

<212> Type : DNA

<211> Length : 33

SequenceName : SEQ ID 119

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 119

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

AVTTDSWGRL H

11

<212> Type : PRT

<211> Length : 11

SequenceName : SEQ ID 120

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gccggttaca ctgacagctg ggggcagctt cat

33

<212> Type : DNA

<211> Length : 33

SequenceName : SEQ ID 121

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SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 121

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

AVTTDSWGQL H

11

<212> Type : PRT

<211> Length : 11

SequenceName : SEQ ID 122

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gccggttaca ctgacagctg ggggaagggtt cat

33

<212> Type : DNA

<211> Length : 33

SequenceName : SEQ ID 123

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 123

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

AVTTDSWGKV H

11

<212> Type : PRT

<211> Length : 11

SequenceName : SEQ ID 124

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

gccggttaca ctgacagctg ggggaagggtg aat

33

<212> Type : DNA

<211> Length : 33

SequenceName : SEQ ID 125

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 125

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

AVTTDSWGKV N

11

<212> Type : PRT

<211> Length : 11

SequenceName : SEQ ID 126

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

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gccggttaca ctgacagctg ggggaagctt ctg 33
<212> Type : DNA
<211> Length : 33
SequenceName : SEQ ID 127
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 127

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AVTTDSWGKL L 11
<212> Type : PRT
<211> Length : 11
SequenceName : SEQ ID 128
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ggaattcatc gatgcagaag gaagtgagc ag 32
<212> Type : DNA
<211> Length : 32
SequenceName : SEQ ID 129
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 129

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gtacacggcc gggtcagggt tctggatata c 31
<212> Type : DNA
<211> Length : 31
SequenceName : SEQ ID 130
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 130

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tctctcatta atgaatgctg gtgtcactca gacccc 36
<212> Type : DNA
<211> Length : 36
SequenceName : SEQ ID 131
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 131

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tagaaaccgg tggccaggca caccagtgtg gc 32
<212> Type : DNA

Final TCR Display PCT.WorkFile.txt

<211> Length : 32
SequenceName : SEQ ID 132
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 132

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ctgctctggt tccgcactc
<212> Type : DNA
<211> Length : 19
SequenceName : SEQ ID 133
SequenceDescription :

19

Custom Codon

Sequence Name : SEQ ID 133

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gagtgcggaa ccagagcag
<212> Type : DNA
<211> Length : 19
SequenceName : SEQ ID 134
SequenceDescription :

19

Custom Codon

Sequence Name : SEQ ID 134

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atatccagaa cccggatcct gccgtgta
<212> Type : DNA
<211> Length : 28
SequenceName : SEQ ID 135
SequenceDescription :

28

Custom Codon

Sequence Name : SEQ ID 135

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tacacggcag gaatccgggt tctggatat
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 136
SequenceDescription :

29

Custom Codon

Sequence Name : SEQ ID 136

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :

Final TCR Display PCT.WorkFile.txt

ggagatatac atatgcagga ggtgacacag 30
<212> Type : DNA
<211> Length : 30
SequenceName : SEQ ID 137
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 137

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tacacggcag gatccgggtt ctggatatt 29
<212> Type : DNA
<211> Length : 29
SequenceName : SEQ ID 138
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 138

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
ggagatatac atatgggtgt cactcagacc 30
<212> Type : DNA
<211> Length : 30
SequenceName : SEQ ID 139
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 139

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cccaagctta gtctgctcta ccccaggcct cggc 34
<212> Type : DNA
<211> Length : 34
SequenceName : SEQ ID 140
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 140

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gccggccatg gccaaacagg aggtgacgca gattcct 37
<212> Type : DNA
<211> Length : 37
SequenceName : SEQ ID 141
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 141

Sequence

Final TCR Display PCT.WorkFile.txt

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cttcttaaag aattcttaat taacctaggt tattaggaac tttctgggct ggggaag 57

<212> Type : DNA
<211> Length : 57
SequenceName : SEQ ID 142
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 142

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tcacagcgcg caggctgggtg tcactcagac cccaaa 36
<212> Type : DNA
<211> Length : 36
SequenceName : SEQ ID 143
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 143

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
cgagagcccg tagaactgga cttg 24
<212> Type : DNA
<211> Length : 24
SequenceName : SEQ ID 144
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 144

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tctctcatta atgaatgctg gtgtcactca gacccc 36
<212> Type : DNA
<211> Length : 36
SequenceName : SEQ ID 145
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 145

Sequence

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
tagaaaccgg tggccaggca caccagtgtg g 31
<212> Type : DNA
<211> Length : 31
SequenceName : SEQ ID 146
SequenceDescription :

Custom Codon

Final TCR Display PCT.WorkFile.txt

Sequence Name : SEQ ID 146

Sequence

```

-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atgcagaagg aagtggagca gaactctgga cccctcagtg ttccagaggg agccattgcc      60
tctctcaact gcacttacag tgaccgaggt tcccagtcct tcttctggta cagacaatat      120
tctgggaaaa gccctgagtt gataatgtcc atatactcca atggtgacaa agaagatgga      180
aggttttacag cacagctcaa taaagccagc cagtatgttt ctctgctcat cagagactcc      240
cagcccagtg attcagccac ctacctctgt gccgttacaa ctgacagctg ggggaaattg      300
cagtttggag cagggacca ggttgtggtc accccagata tccagaacct tgaccctgcc      360
gtgtaccagc tgagagactc taaatccagt gacaagtctg tctgcctatt caccgatttt      420
gattctcaaa caaatgtgtc acaaagtaag gattctgatg tgtatatcac agacaaatgt      480
gtgctagaca tgaggtctat ggacttcaag agcaacagtg ctgtggcctg gagcaacaaa      540
tctgactttg catgtgcaaa cgccttcaac aacagcatta ttccagaaga caccttcttc      600
cccagcccag aaagtctcta a                                621
<212> Type : DNA
<211> Length : 621
      SequenceName : SEQ ID 147
      SequenceDescription :

```

Custom Codon

Sequence Name : SEQ ID 147

Sequence

```

-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atgaacgctg gtgtcactca gaccccaaaa ttccagggtcc tgaagacagg acagagcatg      60
aactgacagt gtgcccagga tatgaaccat gaatacatgt cctgggtatcg acaagaccca      120
ggcatggggc tgaggctgat tcattactca gttgggtgctg gtatcactga ccaaggagaa      180
gtccccaatg gctacaatgt ctccagatca accacagagg atttcccgct caggctgctg      240
tcggctgctc cctcccagac atctgtgtac ttctgtgcc a gcaggccggg actagcggga      300
gggcgaccag agcagtactt cgggcccgggc accagggtca cggtcacaga ggacctgaaa      360
aacgtgttcc caccgaggt cgctgtgttt gagccatcag aagcagagat ctcccacacc      420
caaaaggcca cactggtgtg cctggccaca ggcttctacc ccgaccacgt ggagctgagc      480
tggtgggtga atgggaagga ggtgcacagt ggggtctgca cagaccgca gccctcaag      540
gagcagcccc ccctcaatga ctccagatac gctctgagca gccgcctgag ggtctcggcc      600
accttctggc aggacccccg caaccacttc cgctgtcaag tccagttcta cgggctctcg      660
gagaatgacg agtggacca ggatagggcc aaacccgtca cccagatcgt cagcgccgag      720
gcctggggta gagcagacta a                                741
<212> Type : DNA
<211> Length : 741
      SequenceName : SEQ ID 148

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Final TCR Display PCT.WorkFile.txt
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 148

Sequence

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<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
MQKEVEQNSG PLSVPEGAIA SLNCTYSDRG SQSFFWYRQY SGKSPELIMS IYSNGDKEDG      60
RFTAQLNKAS QYVSLIIRDS QPSDSATYLC AVTTDSWGKL QFGAGTQVVV TPDIQNPDP      120
VYQLRDSKSS DKSVCFLTDF DSQTNVSQSK DSDVYITDKC VLDMRSMDFK SNSAVALWSNK      180
SDFACANAFN NSIIPEDTFF PSPESS      206
<212> Type : PRT
<211> Length : 206
      SequenceName : SEQ ID 149
      SequenceDescription :

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Sequence

```

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
MNAGVTQTPK FQVLKTGQSM TLQCAQDMNH EYMSWYRQDP GMGLRLIHYS VGAGITDQGE      60
VPNGYNVSRs TTEDFPLRLL SAAPSQTSVY FCASRPGLAG GRPEQYFGPG TRLTVTEDLK      120
NVFPPEVAVF EPSEAEISHT QKATLVCLAT GFYPDHVELS WWVNGKEVHS GVCTDPQPLK      180
EQPALNDSRY ALSSRLRVSA TFWQDPRNHF RCQVQFYGLS ENDEWTQDRA KPVTQIVSAE      240
AWGRAD      246
<212> Type : PRT
<211> Length : 246
      SequenceName : SEQ ID 150
      SequenceDescription :

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Sequence

```

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gttaactacg tcaggtggca cttttcgggg aaatgtgcgc ggaacccta tttgtttatt      60
tttctaataa cattcaata tgtatccgct catgagacaa taaccctgat aaatgcttca      120
ataatattga aaaaggaaga gtatgagtat tcaacatttc cgtgtcgccc ttattccctt      180
ttttgcggca ttttgccttc ctgtttttgc tcaccagaa acgctggtga aagtaaaaga      240
tgctgaagat cagttgggtg cagcagtggtg ttacatcgaa ctggatctca acagcggtaa      300
gatccttgag agttttcgcc ccgaagaacg ttctccaatg atgagcactt ttaaagttct      360
gctatgtggc gcggtattat cccgtgttga cgccgggcaa gagcaactcg gtcgccgcat      420
acactattct cagaatgact tggttgagta ctcaccagtc acagaaaagc atcttacgga      480
tgccatgaca gtaagagaat tatgcagtcg tgccataacc atgagtata acactgcggc      540
caacttactt ctgacaacga tcggaggacc gaaggagcta accgcttttt tgcacaacat      600
gggggatcat gtaactcgcc ttgatcggtg ggaaccggag ctgaatgaag ccataccaaa      660
cgacgagcgt gacaccacga tgcctgtagc aatggcaaca acgttgcgca aactattaac      720

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Final TCR Display PCT.WorkFile.txt

tggcgaacta cttactctag cttcccggca acaattaata gactggatgg aggcggataa	780
agttgcagga ccacttctgc gctcggccct tccggctggc tggtttattg ctgataaatc	840
tggagccggt gagcgtgggt ctgcgggtat cattgcagca ctggggccag atggtaagcc	900
ctcccgtatc gtagttatct acacgacggg gagtcaggca actatggatg aacgaaatag	960
acagatcgct gagatagggt cctcactgat taagcattgg taactgtcag accaagttta	1020
ctcatatata ctttagattg atttaccccc gttgataatc agaaaagccc caaaaacagg	1080
aagattgtat aagcaaatat ttaaattgta aacgttaata ttttgttaaa attcgcgtta	1140
aatttttgtt aaatcagctc attttttaac caataggccg aaatcggcaa aatcccttat	1200
aaatcaaaaag aatagcccga gatagggttg agtggtgttc cagtttgga caagagtcca	1260
ctattaaaga acgtggactc caacgtcaaa gggcgaaaaa ccgtctatca gggcgatggc	1320
ccactacgtg aaccatcacc caaatcaagt tttttggggg cgagggtgccg taaagcacta	1380
aatcgggaacc ctaaaggag cccccgattt agagcttgac ggggaaagcg aacgtggcga	1440
gaaaggaagg gaagaaagcg aaaggagcgg gcgctagggc gctggcaagt gtagcggta	1500
cgctgcgcgt aaccaccaca cccgccgcgc ttaatgcgcc gctacagggc gcgtaaaagg	1560
atctagggtg agatcctttt tgataatctc atgacaaaaa tcccttaacg tgagttttcg	1620
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ccaaatactg ttcttctagt gtagccgtag ttaggccacc acttcaagaa ctctgtagca	1860
ccgcctacat acctcgctct gctaatacctg ttaccagtgg ctgctgccag tggcgataag	1920
tcgtgtctta ccgggttgga ctcaagacga tagttaccgg ataaggcgca gcggtcgggc	1980
tgaacggggg gttcgtgcac acagcccagc ttggagcgaa cgacctacac cgaactgaga	2040
tacctacagc gtgagctatg agaaagcgcc acgcttcccg aaggagagaaa ggcggacagg	2100
tatccggtaa gcggcagggt cggaacagga gagcgcacga gggagcttcc agggggaaac	2160
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ctattttcaag gagacagtca taatgaaata cctattgcct acggcagccg ctggattgtt	2520
attactcgcg gccagccgg ccatggcca gaaggaagt gagcagaact ctggaccctt	2580
cagtgttcca gagggagcca ttgcctctct caactgcact tacagtgacc gaggttccca	2640
gtccttcttc tggtagacac aatattctgg gaaaagccct gagttgataa tgtccatata	2700
ctccaatggt gacaaagaag atggaaggtt tacagcacag ctcaataaag ccagccagta	2760

Final TCR Display PCT.WorkFile.txt

tgttttctctg	ctcatcagag	actcccagcc	cagtgtattca	gccacctacc	tctgtgccgt	2820
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tgatgtgtat	atcacagaca	aatgtgtgct	agacatgagg	tctatggact	tcaagagcaa	3060
cagtgtctgtg	gcctggagca	acaaatctga	ctttgcatgt	gcaaacgcct	tcaacaacag	3120
cattattcca	gaagacacct	tcttccccag	cccagaaagt	tcctaataac	ctagggttaat	3180
taagaattct	ttaagaagga	gatatacata	tgaaaaaatt	attattcgca	attccttttag	3240
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tcacggtcac	agaggacctg	aaaaacgtgt	tcccacccga	ggtcgctgtg	tttgagccat	3660
cagaagcaga	gatctcccac	acccaaaagg	ccacactggt	gtgcctggcc	acaggcttct	3720
accccgacca	cgtggagctg	agctgggtggg	tgaatgggaa	ggaggtgcac	agtgggggtct	3780
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gcagccgcct	gaggggtctcg	gccaccttct	ggcaggaccc	ccgcaaccac	ttccgctgtc	3900
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tgtggaatgc	tacaggcggt	gtagtttgta	ctggtgacga	aactcagtgt	tacggtacat	4200
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agaataatag	gttccgaaat	aggcaggggg	cattaactgt	ttatacgggc	actgttactc	4500
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Final TCR Display PCT.WorkFile.txt

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tgagtcgtat tacggactgg ccgtcgtttt acaacgtcgt gactgggaaa accctggcgt	5340
tacccaactt aatcgccttg cagcacatcc ccctttcgcc agctggcgta atagcgaaga	5400
ggcccgacc gatcgccctt cccaacagtt gcgcagcctg aatggcgaat ggcgcttcgc	5460
ttggtaataa agcccgcttc ggcgggcttt ttttt	5495

<212> Type : DNA
 <211> Length : 5495
 SequenceName : SEQ ID 151
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 151

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :

agctgcagct aatacgactc actataggaa caggccacca tgggccagaa ggaagtggag	60
cagaactctg gaccctcag tgttcagag ggagccattg cctctctcaa ctgcacttac	120
agtgaccgag gttcccagtc cttcttctgg tacagacaat attctgggaa aagccctgag	180
ttgataatgt ccatatactc caatgggtgac aaagaagatg gaaggtttac agcacagctc	240
aataaagcca gccagtatgt ttctctgctc atcagagact cccagcccag tgattcagcc	300
acctacctct gtgccgttac aactgacagc tgggggaaat tgcagtttg agcagggacc	360
caggttgtgg tcaccgggtg aggcggttca ggcggagggt gatccggcgg tggcgggtcg	420
aacgctgggtg tctctcagac cccaaaattc caggctctga agacaggaca gagcatgaca	480
ctgcagtgtg cccaggatat gaaccatgaa tacatgtcct ggtatcgaca agaccaggc	540
atggggctga ggctgattca ttactcagtt ggtgctggta tctactgacca aggagaagtc	600
cccaatggct acaatgtctc cagatcaacc acagaggatt tcccgtcag gctgctgtcg	660
gctgctccct cccagacatc tgtgtacttc tgtgccagca ggccgggact agcgggaggg	720
cgaccagagc agtacttcgg gccgggcacc aggtctacgg tcacagagga cctgaaaaac	780
gtgttccac ccgaggctgc tgtgtttgag ccatcagaag cagagatctc ccacacccaa	840
aaggccacac tgggtgtgcct ggccacaggc ttctaccccg accacgtgga gctgagctgg	900
tgggtgaatg ggaaggaggt gcacagtggg gtcagcacag accgcagcc cctcaaggag	960

Final TCR Display PCT.WorkFile.txt

cagccccgcc tcaatgactc cagatacgct ctgagcagcc gcctgagggg ctcggccacc	1020
ttctggcagg acccccgcaa ccacttccgc tgtcaagtcc agttctacgg gctctcggag	1080
aatgacgagt ggacccagga tagggccaaa cccgtcaccc agatcgtcag cgccgaggcc	1140
tggggtagag cagacggtgg aggcggttca ctcagcagca ccctgacgct gagcaaagca	1200
gactacgaga aacacaaagt ctacgcctgc gaagtcaccc atcagggcct gagttcgccc	1260
gtcacaaaga gcttcaaccg cggagagtca taagaattct cag	1303

<212> Type : DNA
 <211> Length : 1303
 SequenceName : SEQ ID 152
 SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 152

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :

MGQKEVEQNS GPLSVPEGAI ASLNCTYS DR GSQSFFWYRQ YSGKSPELIM SIYSNGDKED	60
GRFTAQLNKA SQYVSL LIRD SPSDSATYLC AVTTDSWGKL QFGAGTQVVV TGGGSGGGG	120
SGGGGSNAGV TQTPKFQVLK TGQSM TLQCA QDMNHEYMSW YRQDPGMGLR LIHYSVGAGI	180
TDQGEVPNGY NVSRSTTEDF PLRLLSAAPS QTSVYFCASR PGLAGGRPEQ YFGPGTRLTV	240
TEDLKNVFPF EVAVFEPSEA EISHTQKATL VCLATGFYPD HVELS W WNG KEVHSGVSTD	300
PQPLKEQPAL NDSRYALSSR LRVSATFWQD PRNHFR CQVQ FYGLSENDEW TQDRAK PVTQ	360
IVSAEAWGRA DGGGGSLSST LTLSKADYEK HKVYACEVTH QGLSSPVTKS FNRGES	416

<212> Type : PRT
 <211> Length : 416
 SequenceName : SEQ ID 153
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :

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gaatctgcag tccaccgtga aaagcccctt tacacgcgcc ttggggataa acaaataaaa	120
agatttatgt aagtttatac ataggcgagt actctgttat tgggactatt tacgaagtta	180
ttataacttt ttccttctca tactcataag ttgtaaaggc acagcgggaa taagggaaaa	240
aacgccgtaa aacggaagga caaaaacgag tgggtctttg cgaccacttt cattttctac	300
gacttctagt caaccacgt gctcacccaa tgtagcttga cctagagttg tcgccattct	360
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atacaccgcy ccataatagg gcataactgc ggcccgttct cgttgagcca gcggcgatatg	480
tgataagagt cttactgaac caactcatga gtggtcagtg tcttttcgta gaatgcctac	540
cgtactgtca ttctcttaat acgtcacgac ggtattggta ctcactattg tgacgccggt	600

Final TCR Display PCT.WorkFile.txt

tgaatgaaga ctgttgctag cctcctggct tcctcgattg gcgaaaaaac gtgttggtacc	660
ccctagtaca ttgagcggaa ctagcaaccc ttggcctcga cttacttcgg tatggtttgc	720
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cgacgaacgt ttgtttttt ggtggcgatg gtcgccacca aacaaacggc ctagtctcg	1320
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ccaacctgag ttctgctatc aatggcctat tccgcgtcgc cagcccgact tgcccccaa	1560
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gctaagtaat tacgtcgacc gtgctgtcca aagggtgac ctttcgcccg tctcgcgt	2100
tgcgtaatt aactcaatc gagtgagtaa tccgtggggt ccgaaatgtg aaatacgaag	2160
gccgagcata caacacacct taacactcgc ctattgttaa agtgtgtcct ttgtcgatac	2220
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agggggccat ggctcgagct taagtgaccg gcagcaaaat gttgcagcac tgaccctttt	2340
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cgcggtactac gccataaaag aggaatgctg agacacgcc aagagtgtgg cgtataccac	2520
gtgagagtca tgtagacga gactacggcg tatcaattcg gtcggggctg tgggcgggtg	2580
tgggcgactg cgcgggactg cccgaacaga cgagggccgt aggcgaatgt ctgttcgaca	2640

Final TCR Display PCT.WorkFile.txt

ctggcagagg ccctcgacgt acacagtctc caaaagtggc agtagtggct ttgcgcgct 2699

<212> Type : DNA
 <211> Length : 2699
 SequenceName : SEQ ID 154
 SequenceDescription :

Custom Codon

 Sequence Name : SEQ ID 154

Sequence

 <213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :

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aatattcttg gaaaagccct gagttgataa tgtccatata ctccaatggg gacaaagaag	180
atggaagggt tacagcacag ctcaataaag ccagccagta tgtttctctg ctcatcagag	240
actcccagcc cagtgattca gccacctacc tctgtgccgt tacaactgac agctggggga	300
aattgcagtt tggagcaggg acccaggttg tggtcaccgg tggaggcggg tcaggcggag	360
gtggatccgg cggtagcggg tcgaacgctg gtgtcactca gaccccaaaa ttccaggtcc	420
tgaagacagg acagagcatg aactgcagt gtgcccagga tatgaacat gaatacatgt	480
cctggtatcg acaagacca ggcattggggc tgaggctgat tcattactca gttggtgctg	540
gtatcactga ccaaggagaa gtccccaatg gctacaatgt ctccagatca accacagagg	600
atttcccgt caggctgctg tcggctgctc cctcccagac atctgtgtac ttctgtgcca	660
gcaggccggg actagcggga gggcgaccag agcagtactt cgggccgggc accaggctca	720
cggtcacaga ggacctgaaa aacgtgttcc caccgagggt cgctgtgttt gagccatcag	780
aagcagagat ctccacacc caaaaggcca cactgggtgtg cctggccaca ggcttctacc	840
ccgaccacgt ggagctgagc tgggtgggtga atgggaagga ggtgcacagt ggggtcagca	900
cagaccgca gccctcaag gacgagccc cctcaatga ctccagatac gctctgagca	960
gccgcctgag ggtctcgcc accttctggc aggacccccg caaccacttc cgctgtcaag	1020
tccagttcta cgggctctcg gagaatgacg agtggacca ggatagggcc aaaccctca	1080
cccagatcgt cagcgccgag gcctggggta gagcagacgg tggaggcggg tcaactcagca	1140
gcacctgac gctgagcaaa gcagactacg agaaacacaa agtctacgcc tgcgaagtca	1200
cccatcaggg cctgagttcg cccgtcacia agagcttcaa ccgcggagag tcataagaat	1260
tc	1262

<212> Type : DNA
 <211> Length : 1262
 SequenceName : SEQ ID 155
 SequenceDescription :

Custom Codon

 Sequence Name : SEQ ID 155

Final TCR Display PCT.WorkFile.txt

Sequence

```

-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gctggtgtca ctcagacccc aaaattccag gtcctgaaga caggacagag catgacactg      60
cagtgtgccc aggatatgaa ccatgaatac atgtcctggt atcgacaaga cccaggcatg      120
gggctgaggc tgattcatta ctcagttggt gctggtatca ctgaccaagg agaagtcccc      180
aatggctaca atgtctccag atcaaccaca gaggatttcc cgctcagggt gctgtcggct      240
gctccctccc agacatctgt gtacttctgt gccagcaggc cgggactagc gggaggggtga      300
ccagagcagt acttcggggc gggcaccagg ctcacggtca cagaggacct gaaaaacgtg      360
ttcccacccg aggtcgtgtg gtttgagcca tcagaagcag agatctccca caccctaaag      420
gccacactgg tgtgcctggc cacaggcttc taccgagacc acgtggagct gagctggtgg      480
gtgaatggga aggaggtgca cagtgggggtc tgcacagacc cgcagcccct caaggagcag      540
cccgccctca atgactccag atacgctctg agcagccgcc tgagggtctc ggccaccttc      600
tggcaggacc cccgcaacca cttccgctgt caagtccagt tctacgggct ctcggagaat      660
gacgagtgga cccaggatag ggccaaaccc gtcaccacga tcgtcagcgc cgaggcctgg      720
ggtagagcag ac                                                                732
<212> Type : DNA
<211> Length : 732
      SequenceName : SEQ ID 156
      SequenceDescription :

```

Custom Codon

```

-----
Sequence Name : SEQ ID 156

```

Sequence

```

-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AGVTQTPKFQ VLKTGQSM TL QCAQDMNHEY MSWYRQDPGM GLRLIHYSVG AGITDQGEVP      60
NGYNVSRSTT EDFPLRLLSA APSQTSVYFC ASRPLAGGX PEQYFGPGTR LTVTEDLKNV      120
FPPEVAVFEP SEAEISHTQK ATLVCLATGF YPDHVELSWW VNGKEVHSGV CTDQPPLKEQ      180
PALNDSRYAL SSRLRVSATF WQDPRNHFR C QVQFYGLSEN DEWTQDRAKP VTQIVSAEAW      240
GRAD                                                                244
<212> Type : PRT
<211> Length : 244
      SequenceName : SEQ ID 157
      SequenceDescription :

```

Feature

```

-----
Sequence: SEQ ID 157:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
<222> LocationTo : 244
      Other Information : X denotes an amino acid residue encoded by an opal
codon, this is normally translated as a tryptophan
      CDSJoin : No

```

Sequence

```

-----

```

Final TCR Display PCT.WorkFile.txt

```

<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gctggtgtca ctcagacccc aaaattccag gtcctgaaga caggacagag catgacactg      60
tagtgtgccc aggatatgaa ccatgaatac atgtcctggt atcgacaaga cccaggcatg      120
gggctgaggc tgattcatta ctcagttggt gctggtatca ctgaccaagg agaagtcccc      180
aatggctaca atgtctccag atcaaccaca gaggatttcc cgctcaggct gctgtcggct      240
gctccctccc agacatctgt gtacttctgt gccagcaggc cgggactagc gggagggcga      300
ccagagcagt acttcgggcc gggcaccagg ctcacggtca cagaggacct gaaaaacgtg      360
ttcccacccg aggtcgctgt gtttgagcca tcagaagcag agatctccca caccctaaag      420
gccacactgg tgtgcctggc cacaggcttc taccctgacc acgtggagct gagctggtgg      480
gtgaatggga aggaggtgca cagtgggggtc tgcacagacc cgcagcccct caaggagcag      540
cccgccctca atgactccag atacgctctg agcagccgcc tgagggtctc ggccaccttc      600
tggcaggacc cccgcaacca cttccgctgt caagtccagt tctacgggct ctcggagaat      660
gacgagtgga cccaggatag ggccaaaccc gtcaccaga tcgtcagcgc cgaggcctgg      720
ggtagagcag ac                                     732
<212> Type : DNA
<211> Length : 732
      SequenceName : SEQ ID 158
      SequenceDescription :

```

Custom Codon

```

-----
Sequence Name : SEQ ID 158

```

Sequence

```

-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
gctggtgtca ctcagacccc aaaattccag gtcctgaaga caggacagag catgacactg      60
cagtgtgccc aggatatgaa ccatgaatac atgtcctggt atcgacaaga cccaggcatg      120
gggctgaggc tgattcatta ctcagttggt gctggtatca ctgaccaagg agaagtcccc      180
aatggctaca atgtctccag atcaaccaca gaggatttcc cgctcaggct gctgtcggct      240
gctccctccc agacatctgt gtacttctgt gcctcaggc cggggctgat gagtgcggaa      300
ccagagcagt acttcgggcc gggcaccagg ctcacggtca cagaggacct gaaaaacgtg      360
ttcccacccg aggtcgctgt gtttgagcca tcagaagcag agatctccca caccctaaag      420
gccacactgg tgtgcctggc caccggtttc taccctgacc acgtggagct gagctggtgg      480
gtgaatggga aggaggtgca cagtgggggtc tgcacagacc cgcagcccct caaggagcag      540
cccgccctca atgactccag atacgctctg agcagccgcc tgagggtctc ggccaccttc      600
tggcaggacc cccgcaacca cttccgctgt caagtccagt tctacgggct ctcggagaat      660
gacgagtgga cccaggatag ggccaaaccc gtcaccaga tcgtcagcgc cgaggcctgg      720
ggtagagcag actaagcttg aattc                                     745
<212> Type : DNA
<211> Length : 745
      SequenceName : SEQ ID 159

```

Final TCR Display PCT.WorkFile.txt
SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 159

Sequence

```

-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
MNAGVTQTPK FQVLKTGQSM TLQCAQDMNH EYMSWYRQDP GMGLRLIHYS VGAGITDQGE      60
VPNGYNVSRS TTEDFPLRLL SAAPSQTSVY FCASRPGLMS AEPEQYFGPG TRLTVTEDLK      120
NVFPPEVAVF EPSEAEISHT QKATLVCLAT GFYPDHVELS WWVNGKEVHS GVCTDPQPLK      180
EQPALNDSRY ALSSRLRVSA TFWQDPRNHF RCQVQFYGLS ENDEWTQDRA KPVTQIVSAE      240
AWGRAD                                             246
<212> Type : PRT
<211> Length : 246
      SequenceName : SEQ ID 160
      SequenceDescription :

```

Sequence

```

-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
AGVTQTPKFQ VLKTGQSMTL QCAQDMNHEY MSWYRQDPGM GLRLIHYSVG AGITDQGEVP      60
NGYNVSRSTT EDFPLRLLSA APSQTSVYFC ASRPGLMSAQ PEQYFGPGTR LTVTEDLKNV      120
FPPEVAVFEP SEAEISHTQK ATLVCLATGF YPDHVELSWW VNGKEVHSGV CTDQPQLKEQ      180
PALNDSRYAL SSRLRVSATF WQDPRNHFRC QVQFYGLSEN DEWTQDRAKP VTQIVSAEAW      240
GRAD                                             244
<212> Type : PRT
<211> Length : 244
      SequenceName : SEQ ID 161
      SequenceDescription :

```

Sequence

```

-----
<213> OrganismName : Synthetic Sequence
<400> PreSequenceString :
atgcaggagg tgacacagat tcctgcagct ctgagtgtcc cagaaggaga aaacttggtt      60
ctcaactgca gtttactga tagcgctatt tacaacctcc agtggttttag gcaggaccct      120
gggaaaggctc tcacatctct gttgcttatt cagtcaagtc agagagagca aacaagtgga      180
agacttaatg cctcgctgga taaatcatca ggacgtagta ctttatacat tgcagcttct      240
cagcctgggtg actcagccac ctacctctgt gctgtgaggc ccacatcagg aggaagctac      300
atactacat ttggaagagg aaccagcctt attgttcatc cgtatatcca gaaccctgac      360
cctgccgtgt accagctgag agactctaaa tccagtgaca agtctgtctg cctattcacc      420
gattttgatt ctcaaacaaa tgtgtcacia agtaaggatt ctgatgtgta tatcacagac      480
aaatgtgtgc tagacatgag gtctatggac ttcaagagca acagtgtgtg ggcctggagc      540
aacaaatctg actttgcatg tgcaaacgcc ttcaacaaca gcattattcc agaagacacc      600
ttcttcccca gccagaaaag ttcctaa                                             627
<212> Type : DNA

```

Final TCR Display PCT.WorkFile.txt

<211> Length : 627
 SequenceName : SEQ ID 162
 SequenceDescription :

Custom Codon

 Sequence Name : SEQ ID 162

Sequence

 <213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 atgggtgtca ctcagacccc aaaattccag gtcctgaaga caggacagag catgacactg 60
 cagtgtgccc aggatatgaa ccatgaatac atgtcctggt atcgacaaga cccaggcatg 120
 gggctgaggc tgattcatta ctcagttggt gctggtatca ctgaccaagg agaagtcccc 180
 aatggctaca atgtctccag atcaaccaca gaggatttcc cgctcaggct gctgtcggct 240
 gtcctctccc agacatctgt gtacttctgt gccagcagtt acgtcgggaa caccggggag 300
 ctgttttttg gagaaggctc taggctgacc gtactggagg acctgaaaaa cgtgttccca 360
 cccgaggctg ctgtgtttga gccatcagaa gcagagatct cccacacca aaaggccaca 420
 ctggtgtgcc tggccacagg cttctacccc gaccacgtgg agctgagctg gtgggtgaat 480
 gggaaggagg tgcacagtgg ggtctgcaca gaccgcagc ccctcaagga gcagcccgcc 540
 ctcaatgact ccagatacgc tctgagcagc cgcctgaggg tctcgccac cttctggcag 600
 gacccccgca accacttccg ctgtcaagtc cagtctacg ggctctcgga gaatgacgag 660
 tggacccagg atagggccaa acccgtcacc cagatcgta gcgccgaggc ctggggtaga 720
 gcagactaa 729

<212> Type : DNA

<211> Length : 729
 SequenceName : SEQ ID 163
 SequenceDescription :

Custom Codon

 Sequence Name : SEQ ID 163

Sequence

 <213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 MQEVTQIPAA LSVPEGENLV LNCSFTDSAI YNLQWFRQDP GKGLTSLLLI QSSQREQTSG 60
 RLNASLDKSS GRSTLYIAAS QPGDSATYLC AVRPTSGGSY IPTFGRGTSL IVHPYIQNPD 120
 PAVYQLRDSK SSDKSVCLFT DFDSQTNVSQ SKDSDVYITD KCVLDMRSMF FKSNSAVAWS 180
 NKSDFACANA FNNSIIPEDT FFPSPESS 208

<212> Type : PRT

<211> Length : 208
 SequenceName : SEQ ID 164
 SequenceDescription :

Sequence

 <213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :
 MGVTTQPKFQ VLKGTQSMFL QCAQDMNHEY MSWYRQDPGM GLRLIHYSVG AGITDQGEVP 60

Final TCR Display PCT.WorkFile.txt

NGYNVSRSTT EDFPLRLLSA APSQTSVYFC ASSYVGNTGE LFFGEGSRLT VLEDLKNVFP	120
PEVAVFEPSE AEISHTQKAT LVCLATGFYP DHVELSWWVN GKEVHSGVCT DPQPLKEQPA	180
LNDSTRYALSS RLRVSATFWQ DPRNHFRQV QFYGLSENDE WTQDRAKPVV QIVSAEAWGR	240
AD	242

<212> Type : PRT
 <211> Length : 242
 SequenceName : SEQ ID 165
 SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
 <400> PreSequenceString :

ttcctggcct tttgctggcc ttttgctcac atgtaatgtg agttagctca aaggaccgga	60
aaacgaccgg aaaacgagtg tacattacac tcaatcgagt ctcataggc accccaggct	120
ttacacttta tgcttccggc tcgtatgttg gagtaatccg tgggggccga aatgtgaaat	180
acgaaggccg agcatacaac tgtggaattg tgagcggata acaatttcac acaggaaaca	240
gctatgacca acaccttaac actcgccat tgttaaagtg tgcctttgt cgatactggt	300
tgattacgcc aagctacgta cttagtatt ctatttcaag gagacagtca actaatgcgg	360
ttcgatgcat gaattcataa gataaagttc ctctgtcagt taatgaaata cctattgcct	420
acggcagccg ctggattgtt attactcgcg attactttat ggataacgga tgccgtcggc	480
gacctaaca taatgagcgc gccagccgg ccatggccaa acaggagggtg acgcagattc	540
ctgcagctct cgggtcggcc ggtaccggtt tgcctccac tgcgtctaag gacgtcgaga	600
gagtgtccca gaaggagaaa acttggttct caactgcagt ttcactgata ctacagggt	660
cttctcttt tgaaccaaga gttgacgtca aagtgactat gcgctattta caacctccag	720
tggtttaggc aggaccctgg gaaaggcttc cgcgataaat gttggagggtc accaaatccg	780
tcctgggacc ctttccagag acatctctgt tgcttattca gtcaagtcag agagagcaaa	840
caagtggaag tgtagagaca acgaataagt cagttcagtc tctctcgttt gttcaccttc	900
acttaatgcc tcgctggata aatcatcagg acgtagtact ttatacattg tgaattacgg	960
agcgacctat ttagtagtcc tgcacatga aatatgtaac cagcttctca gcctggtgac	1020
tcagccacct acctctgtgc tgtgaggccc gtcgaagagt cggaccactg agtcggtgga	1080
tggagacacg aactccggg acatcaggag gaagctacat acctacattt ggaagaggaa	1140
ccagccttat ttagtcctc cttcgatgta tggatgtaaa cttctcctt ggtcgggaata	1200
tgttcatccg tatatccaga acccgatcc tgccgtgtac cagctgagag acaagtaggc	1260
atataggtct tgggcctagg acggcacatg gtcgactctc actctaaatc cagtgacaag	1320
tctgtctgcc tattcaccga ttttgattct tgagatttag gtcactgttc agacagacgg	1380
ataagtggct aaaactaaga caaacaatg tgtcacaaag taaggattct gatgtgtata	1440
tcacagacaa gtttggtttac acagtgtttc attcctaaga ctacacatat agtgtctgtt	1500
atgtgtgcta gacatgaggt ctatggactt caagagcaac agtgctgtgg tacacacgat	1560

Final TCR Display PCT.WorkFile.txt

ctgtactcca gatacctgaa gttctcgttg tcacgacacc cctggagcaa caaatctgac	1620
tttgcattgtg caaacgcctt caacaacagc ggacctcgtt gtttagactg aaacgtacac	1680
gtttgcggaa gttgttgtcg attattccag aagacacctt cttccccagc ccagaaagtt	1740
cctaataacc taataagggtc ttctgtggaa gaaggggtcg ggtctttcaa ggattattgg	1800
taggttaatt aagaattctt taagaagggg atatacatat gaaaaaatta atccaattaa	1860
ttcttaagaa attcttcccc tatatgtata cttttttaat ttattcgcaa ttcctttagt	1920
tgttcctttc tattctcaca gcgcgcaggc aataagcggt aaggaaatca acaaggaaag	1980
ataagagtgt cgcgcgtccg tggtgtcact cagaccccaa aattccagggt cctgaagaca	2040
ggacagagca accacagtga gtctgggggt ttaagggtcca ggacttctgt cctgtctcgt	2100
tgacactgca gtgtgcccag gatatgaacc atgaatacat gtcctggtat actgtgacgt	2160
cacacgggtc ctatacttgg tacttatgta caggaccata cgacaagacc caggcatggg	2220
gctgaggctg attcattact cagttggtgc gctgttctgg gtccgtaccc cgactccgac	2280
taagtaatga gtcaaccacg tggtatcact gaccaaggag aagtcccca tggctacaat	2340
gtctccagat accatagtga ctgggtcctc ttcaggggtt accgatgtta cagagggtcta	2400
caaccacaga ggatttcccg ctcaggctgc tgtcggctgc tccctcccag gttggtgtct	2460
cctaaagggc gagtccgacg acagccgacg agggaggggt acatctgtgt acttctgtgc	2520
cagcagttac gtcgggaaca ccggggagct tgtagacaca tgaagacacg gtcgtcaatg	2580
cagcccttgt ggccctcga gttttttgga gaaggctcta ggctgaccgt actggaggac	2640
ctgaaaaacg caaaaaacct cttccgagat ccgactggca tgacctctg gactttttgc	2700
tgttcccacc cgaggctcgt gtgtttgagc catcagaagc agagatctcc acaagggtgg	2760
gctccagcga cacaaactcg gtagtcttcg tctctagagg cacacccaaa aggccacact	2820
gggtgtgcctg gccacaggct tctaccccgga gtgtgggttt tccggtgtga ccacacggac	2880
cggtgtccga agatggggct ccacgtggag ctgagctggt ggggtgaatgg gaaggagggtg	2940
cacagtgggg ggtgcacctc gactcgacca cccacttacc cttcctccac gtgtcacccc	3000
tctgcacaga cccgcagccc ctcaaggagc agcccgccct caatgactcc agacgtgtct	3060
gggcgtcggg gagttcctcg tcgggcggga gttactgagg agatacgctc tgagcagccg	3120
cctgagggtc tcggccacct tctggcagga tctatgcgag actcgtcggc ggactcccag	3180
agccggtgga agaccgtcct ccccgcaac cacttccgct gtcaagtcca gttctacggg	3240
ctctcggaga gggggcggtg gtgaaggcga cagttcagggt caagatgccc gagagcctct	3300
atgacgagtg gaccaggat agggccaaac ccgtcaccca gatcgtcagc tactgctcac	3360
ctgggtccta tcccggtttg ggcagtgggt ctagcagtcg gccgaggcct ggggtagagc	3420
agacgcggcc gcacggctcc ggaccccatc tcgtctgcgc cggcgt	3466

<212> Type : DNA
 <211> Length : 3466
 SequenceName : SEQ ID 166
 SequenceDescription :

Final TCR Display PCT.WorkFile.txt

Custom Codon

Sequence Name : SEQ ID 166

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

KYLLPTAAAG LLLLAAPAM AKQEVTPQA ALSVPEGENL VLNCSFTDSA IYNLQWFRQD	60
PGKGLTSLLL IQSSQREQTS GRLNASLDKS SGRSTLYIAA SQPGDSATYL CAVRPTSGGS	120
YIPTFGRGTS LIVHPYIQNP DPAVYQLRDS KSSDKSVCLF TDFDSQTNVS QSKDSDVYIT	180
DKCVLDMRSM DFKSNSAVAW SNKSDFACAN AFNNSIIPED TFFPSPESSM KKLLFAIPLV	240
VPFYSHSAQA GVTQTPKFQV LKTGQSMTLQ CAQDMNHEYM SWYRQDPGMG LRLIHYSVGA	300
GITDQGEVNP GYNVSRSTTE DFPLRLLSAA PSQTSVYFCA SSVVGNTGEL FFGEGSRLTV	360
LEDLKNVFPP EVAVFEPSEA EISHTQKATL VCLATGFYPD HVELSWVNG KEVHSGVCTD	420
PQPLKEQPAL NDSRYALSSR LRVSATFWQD PRNHFRQVQ FYGLSENDEW TQDRAKPVTQ	480
IVSAEAWGRA DAAA	494

<212> Type : PRT

<211> Length : 494

SequenceName : SEQ ID 167

SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

ggatccatgg ccataagtgg agtccctgtg ctaggatttt tcatcatagc tgtgctgatg	60
agcgctcagg aatcatgggc tatcaaagaa gaacatgtga tcatccaggc cgagttctat	120
ctgaatcctg accaatcagg cgagtttatg tttgactttg atgggtgatga gattttccat	180
gtggatatgg caaagaagga gacgggtctgg cggcttgaag aatttgagcg atttgccagg	240
tttgaggctc aagggtgcatt ggccaacata gctgtggaca aagccaacct ggaaatcatg	300
acaaagcgct ccaactatac tccgatcacc aatgtacctc cagaggtaac tgtgctcacg	360
aacagccctg tggaactgag agagcccaac gtcctcatct gtttcatcga caagttcacc	420
ccaccagtgg tcaatgtcac gtggcttcga aatggaaaac ctgtcaccac aggagtgtca	480
gagacagtct tcctgcccag ggaagaccac cttttccgca agttccacta tctccccttc	540
ctgccctcaa ctgaggacgt ttacgactgc aggggtggagc actggggcctt ggatgagcct	600
cttctcaagc actgggagtt tgatgctcca agccctctcc cagagactac agagaacgtg	660
gatgggggtc tgactgatac actccaagcg gagacagatc aacttgaaga caagaagtct	720
gcgttgacga ccgagattgc caatctactg aaagagaagg aaaaactaga gttcatcctg	780
gcagcttacg gatctggtgg tggtctgaac gatatttttg aagctcagaa aatcgaatgg	840
catgagtagg atcc	854

<212> Type : DNA

<211> Length : 854

SequenceName : SEQ ID 168

SequenceDescription :

Final TCR Display PCT.workFile.txt

Custom Codon

Sequence Name : SEQ ID 168

Sequence

<213> OrganismName : Synthetic Sequence

<400> PreSequenceString :

```

aattctactc gtaaagcgag ttgaaggatc atatttagtt gcgtttatga gataagattg      60
aaagcacgtg taaaatgttt cccgcgcgtt ggcacaacta ttacaatgc ggccaagtta      120
taaaagattc taatctgata tgttttaaaa cacctttgcg gcccgagttg tttgcgtacg      180
tgactagcga agaagatgtg tggaccgcag aacagatagt aaaacaaaac ctagtattg      240
gagcaataat cgatttaacc aacacgtcta aatattatga tgggtgtgcat tttttgcggg      300
cgggcctgtt atacaaaaaa attcaagtac ctggccagac tttgccgcct gaaagcatag      360
ttcaagaatt tattgacacg gtaaaagaat ttacagaaaa gtgtcccggc atgttggtgg      420
gcgtgcactg cacacacggt attaatcgca ccggttacat ggtgtgcaga tatttaatgc      480
acaccctggg tattgcgccg caggaagcca tagatagatt cgaaaaagcc agaggtcaca      540
aaattgaaag acaaaattac gttcaagatt tattaattta attaataatta tttgcattct      600
ttaacaaata ctttatccta ttttcaaatt gttgcgcttc ttccagcgaa caaaactat      660
gcttcgcttg ctccgttttag cttgtagccg atcagtggcg ttgttccaat cgacggtagg      720
attaggcccg atattctcca ccacaatgtt ggcaacgttg atgttacgtt tatgcttttg      780
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Final TCR Display PCT.WorkFile.txt

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Final TCR Display PCT.WorkFile.txt

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Final TCR Display PCT.WorkFile.txt

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Final TCR Display PCT.WorkFile.txt

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<211> Length : 10096

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SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 169

Sequence

<213> OrganismName : Synthetic Sequence

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<212> Type : DNA

<211> Length : 941

SequenceName : SEQ ID 170

SequenceDescription :

Custom Codon

Sequence Name : SEQ ID 170

Sequence

<213> OrganismName : Synthetic Sequence

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<212> Type : PRT

Final TCR Display PCT.WorkFile.txt

<211> Length : 116
SequenceName : SEQ ID 171
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
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<212> Type : PRT
<211> Length : 116
SequenceName : SEQ ID 172
SequenceDescription :

Feature

Sequence: SEQ ID 172:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 1
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Other Information : where x is e, q or r
CDSJoin : No

Sequence

<213> OrganismName : Synthetic Sequence
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Sequence

<213> OrganismName : Synthetic Sequence
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SequenceName : SEQ ID 174
SequenceDescription :

Sequence

<213> OrganismName : Synthetic Sequence
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<212> Type : PRT
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SequenceDescription :

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